## -continued ( x i ) SEQUENCE DESCRIPTION: SEQ ID NO:1: ACUCUCUUCC OCAUCOCUOU CUOCOAGOGC CAOCUGUUDO GCUCGCOGUU GAGGACAAAC DCAACGCOOR CARRESON CACAROOPIC CONVECCOR COCCACCOV VCONVCACCO 120 CCACCOAGOO ACCUGAGGA GUCÓGCAUCO ACCOGAUCO AAAACCUCUC GAGAAAGGC UCUAACCAGU CACAGUCGCA 200 ( 2 ) INFORMATION FOR SEQ ID NO:2: ( i ) SEQUENCE CHARACTERISTICS: ( A ) LENGTH: 33 base pairs ( B ) TYPE: sucleic seid ( C ) STRANDEDNESS: single ( D ) TOPOLOGY: Esset (ii) MOLECULE TYPE: mRNA ( $\mathbf{x}$ $\mathbf{i}$ ) SEQUENCE DESCRIPTION: SEQ ID NO2: ACUCUCUUCC OCAUCOCUOU CUOCOAGGGC CAG 3 3 ( 2 ) INFORMATION FOR SEQ ID NOS: ( i ) SEQUENCE CHARACTERISTICS: (A) LENGTH: 12 been pairs ( B ) TYPE: moleic seid ( C ) STRANDEDNESS: single ( D ) TOPOLOGY: Honer ( i i ) MOLECULE TYPE: DNA (genomic) ( x i ) SEQUENCE DESCRIPTION: SEQ ID NO:4: AGCTTIGATC AG 1 2 ( 2 ) INFORMATION FOR SBQ ID NO:4: ( i ) SEQUENCE CHARACTERISTICS: ( A ) LENGTEE 12 base pairs ( B ) TYPE; meleic acid ( C ) STRANDEDNESS: single ( D ) TOPOLOGY: linear ( i i ) MOLECULE TYPE: DNA (genomic) ( x i ) SEQUENCE DESCRIPTION: SEQ ID NO:4: GCACCIGÁIC AA ( 2 ) INFORMATION FOR SEQ ID NO:5: ( i ) SEQUENCE CHARACTERISTICS: ( A ) LENGTH: 8 base pairs ( B ) TYPE: moclein acid ( C ) STRANDEDNESS: single ( D ) TOPOLOGY: linear ( i i ) MOLECULE TYPE: DNA (genomie) ( $\mathbf{z} \cdot \mathbf{i}$ ) SEQUENCE DESCRIPTION: SEQ ID NO.5: GTGATCAA ( 2 ) INFORMATION FOR SEQ ID NO::

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 16 bee pairs
(B) TYPE: mucleio scid
(C) STRANDEUNESS: single
(D) TOPOLOGY; Heest

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| ( i i ) MOLECULE TYPE: DNA (peromic)                             |               |            |            |       |
|--|---------------|------------|------------|-------|
| ( x i ) SEQUENCE DESCRIPTION: SEQ ID NO.                         | •             |            |            | •     |
| GATCTTGATC ACTGCA  |               |            |            | 1.6   |
|  | •             |            |            |       |
| (2) INFORMATION FOR SEQ ID NOUT                                  |               |            |            |       |
| ( i ) SEQUENCE CHARACTERISTICS:                                  |               |            |            |       |
| ( A ) LENGTH: 8 base pass  | •             | -          |            |       |
| (B) TYPE: mekic acid   |               |            |            |       |
| (C) STRANDEDNESS: single   |               | •          |            |       |
| (D) TOPOLOGY: House  |               |            |            |       |
| ( i i ) MOLECULE TYPE: DNA (genomic)                             |               |            |            |       |
| ( $\mathbf{z}$ $\mathbf{i}$ ) SEQUENCE DESCRIPTION: SEQ ID NO:7: |               |            |            |       |
| COGATCCO   |               |            |            | . 1   |
|  |               |            |            |       |
| (2) INFORMATION FOR SEQ ID NO:8:                                 |               | •          |            |       |
|  |               |            | •          |       |
| ( 1 ) SEQUENCE CHARACTERISTICS:                                  | •             |            |            |       |
| ( A ) LENGIH: 8 base pairs                                       |               |            |            |       |
| (B)TYPE: meleic acid.  | •             |            |            |       |
| (C) STRANDEONESS: single<br>(D) TOPCELOGY: linear                | •             |            |            |       |
| ( i i ) MOLECULE TYPE: DNA (genomic)                             |               |            | _          |       |
| ( x i ) SEQUENCE DESCRIPTION: SEQ ID NO:4:                       |               |            |            |       |
|  |               |            |            | •     |
| COGATCCG   | ·             |            |            |       |
| (2) INFORMATION FOR SEQ ID NO-9:                                 |               | •          |            |       |
| ( i ) SEQUENCE CHARACTERISTICS:                                  | •             |            |            | ,     |
| (A) LENGIH: 287 base pairs                                       |               |            | •          |       |
| (B) TYPE: meleia acid  |               |            |            |       |
| (C) STRANDEDNESS: might  |               |            |            |       |
| ( D ) TOPOLOGY: Harm   | •             | •          |            |       |
| - ( i i ) MOLECULE TYPE: DNA (genomic)                           | ,             |            |            |       |
| ( x i ) SEQUENCE DESCRIPTION: SEQ ID NO:9:                       |               | ·          |            |       |
| AATICACOCT GTGGTGTTAT GGTCGGTGG                                  |               |            |            | . 60  |
| ACCOTOCACC AATCCTTCTC GCCTCAGGC                                  |               |            |            | 120   |
| GGTCGTATAA TCACCGCATA ATTCGAGTC                                  |               |            |            | 240   |
| TITITOCICC GACATCATAA CGGTTCCGG                                  |               |            | TTGACAATIA | 287   |
| ATCATCGAAC TAGTTAACTA GTACGCAAG                                  | T TCTCGTAAAA  | AGGGTAT    |            |       |
| ( 2 ) INFORMATION FOR SBQ ID NO:10:                              |               |            |            |       |
| ( i ) SEQUENCE CHARACTERISTICS:                                  | •             | •          |            |       |
| (A) LENGTH: 285 base pairs                                       |               | •          |            |       |
| ( B ) TYPE: melsis seid  | •             |            |            |       |
| (C) STRANDEDNESS: single   |               |            |            |       |
| ( D ) TOPOLOGY: Hear   | •             |            |            | ,     |
| ( i i ) MOLECULE TYPE: DNA (genomic)                             |               |            |            |       |
| ( x i ) SEQUENCE DESCRIPTION: SEQ ID NO:10:                      |               |            |            | 6 0   |
| COATACCCTT TITACOADAA CTIGCGTAC                                  | T AGITAACTAG  | TICOATOATT | AATTOTCAAC | 120   |
| ACCTEATITE AGAATATITG CCGGAACEG                                  | TATGATGTCG    | GAGCAAAAA  | CAITATCCOO | 1 8 0 |
| AACOGGAGTG CGCCTTGAGC GACTCGAAT                                  |               |            |            | 240   |
| ACCACAGETT CCGATTOOCT GCCTGACGC                                  | EC AGAAGCATTG | GIGCACCGIG | CAGTCUAGAI | 1-0   |

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| OCCCOTCOGC ACCCTACCGA CCACCGACC                      | A TAACACCACA GCGTG |    | 2 8 5            |
|--|--------------------|----|------------------|
| ( 2 ) INPORMATION FOR SEQ ID NO:11:                  | ,                  |    |                  |
|  |                    |    |                  |
| ( i ) SEQUENCE CHARACTERISTICS:                      | •                  |    |                  |
| ( A ) LENGTH: 8 base pairs                           |                    |    |                  |
| ( B ) TYPE: models said                              | • •                |    |                  |
| ( C ) STRANDEDNESS: single<br>( D ) TOPOLOGY: Hans:  |                    |    |                  |
| ( i i ) MOLECULE TYPE: DNA (graceria)                | •                  |    |                  |
| (x i ) SEQUENCE DESCRIPTION: SEQ ID NO:11:           |                    |    |                  |
| DOTATAS  |                    |    | •                |
| (2) INFORMATION FOR SEQ ID NO:12:                    |                    |    |                  |
| · · · · · · · · · · · · · · · · · · ·                |                    |    |                  |
| ( i ) SEQUENCE CHARACTERISTICS:                      |                    |    |                  |
| (A) LENGIE: 8 base pairs<br>(B) TYPE: michie ackl    |                    | •  |                  |
| (C) STRANDEDNESS: made                               |                    |    |                  |
| ( D ) TOPOLOGY: Home                                 |                    |    |                  |
| ( 1 i ) MOLECULE TYPE: DNA (genomic)                 |                    |    |                  |
| ( x i ) SEQUENCE DESCRIPTION: SEQ ID NO:12:          |                    |    | . •              |
| CCATATGG   |                    |    | •                |
| ( 2 ) INFORMATION FOR SEQ ID NO:13:                  |                    |    |                  |
| ( 1 ) SEQUENCE CHARACTERISTICS:                      |                    |    |                  |
| ( A ) LENGTH: 8 beso poins                           |                    |    |                  |
| (B) TIPS: meleic acid                                |                    |    | •                |
| ( C ) STRANDEDNESS: single<br>( D ) TOPOLOGY: linear |                    | ·· |                  |
| ( i i ) MOLECULE TYPE: DNA (genomic)                 | •                  |    |                  |
| ( x i ) SEQUENCE DESCRIPTION: SEQ ID NO:13:          | ·                  |    |                  |
| COTTAACO   | ,                  | •  | •                |
| ( 2 ) INFORMATION FOR SEQ ID NO:14:                  |                    |    | -                |
| ( i ) SEQUENCE CHARACTERISTICS:                      |                    |    |                  |
| ( A ) LENGTH: 8 base pairs                           |                    |    |                  |
| (B) TYPE: meleic acid                                |                    |    |                  |
| ( C ) STRANDEDNESS: single<br>( D ) TOPOLOGY: linear |                    |    |                  |
| ( i i ) MOLECULE TYPE: DNA (geocomic)                | •                  |    |                  |
| ( x i ) SEQUENCE DESCRIPTION: SEQ ID NO:144          |                    | :  | <del>-</del> . · |
| CGTTAACG   | •                  | ,  | 8                |
| ( 2 ) Information for SEQ ID No:15:                  |                    |    |                  |
|  |                    |    |                  |
| ( i ) SEQUENCE CHARACTERISTICS:                      |                    |    |                  |
| ( A ) LENGTH: 36 bus pain                            |                    |    |                  |
| (B) TYPE: modeic acid<br>(C) STRANDEDNESS: single    |                    |    |                  |
| ( D ) TOPCLOOY: Head                                 |                    |    |                  |
| ( i i ) MOLECULE TYPE: DNA (genomic)                 | •                  |    |                  |
| ( z i ) SEQUENCE DESCRIPTION: SEQ ID NO:15:          |                    |    |                  |
| GGGAAGTGCT GTGAAATATC CACCTGCGG                      | C CTGAGA           |    | 3 6              |
|  |                    |    |                  |

( 2 ) INFORMATION FOR SEQ ID NO:16:

-continued ( i ) SEQUENCE CHARACTERISTICS: ( A ) LENGTH: 46 base pains ( B ) TYPE: sucloic said (C) STRANDEDNESS: single (D) TOPOLOGY: Illust ( i i ) MOLECULE TYPE: DNA (genomic) ( z i ) SEQUENCE DESCRIPTION: SEQ ID NO:16: CTAGAGGGTA TTAATAATGT ATCGATTTAA ATAAGGAGGA ATAACA ( 2 ) INFORMATION FOR SEQ ID NO:17: ( 1 ) SEQUENCE CHARACTERISTICS: ( A ) LENGTH: 44 base point ( B ) TYPE: modeic soid ( C ) STRANDEDNESS: single ( D ) TOPOLOGY: linear ( i i ) MOLECULE TYPE: DNA (genomic) ( x i ) SEQUENCE DESCRIPTION: SEQ ID NO:17: TATGITATIC CTCCTTATTI AAATCGATAC ATTATTAATA CCCT 4 4 ( 2 ) INFORMATION FOR SEQ ID NO:18: ( i ) SEQUENCE CHARACTERISTICS: ( A ) LENGTE: 22 bess pairs ( B ) TYPE: moleic scid ( C ) STRANDEDNESS: single ( D ) TOPOLOGY: Esser ( i i ) MOLECULE TYPE: DNA (genomic) ( x i ) SEQUENCE DESCRIPTION: SEQ ID NO:18: GATCIATIAA CICAATCIAG AC 2 2 (2) INFORMATION FOR SEQ ID NO:19: ( i ) SEQUENCE CHARACTERISTICS: (A.) LENOTH: 22 base pairs (B) TYPE: mucleic acid (C) STRANDEDNESS: meda ( D ) TOPOLOGY: Henr ( i i ) MOLECULE TYPE: DNA (genomic) ( x i ) SEQUENCE DESCRIPTION: SEQ ID NO:19: TCGAGICTAG ATTGAGTTAA TA 22. ( 2 ) INFORMATION FOR SEQ ID NO:20: ( i ) SEQUENCE CHARACTERISTICS: ( A ) LENGTH: 872 base pairs ( B ) TYPE: sucleic acid ( C ) STRANDEDNESS: single ( D ) TOPOLOGY: finese . ( i i ) MOLECULE TYPE: DNA (genomic) ( x 1 ) SEQUENCE DESCRIPTION: SEQ ID NO:20: AAGCTITTCT CATTAAGGGA AGATTTCCCC AGGCAGCTCT TICAAGGCCT AAAAGGTCCA 60 TOAGCTCCAT GGATTCTTCC CTGTTAAGAA CTTTATCCAT TTTTGCAAAA ATTGCAAAAG ANTAGOGATT TECECAAATA GTTTTGETAG GEETCAGAAA AAGEETCEAE ACCETTACTA 1 8 0 CTTGAGAGAA AGGGTGGAGO CAGAGGCGGC CTCGGCCTCT TATATATTAT AAAAAAAAG 240

GCCACAGGA GGAGCTGCTT ACCCATGGAA TGCAGCCAAA CCATGACCTC AGGAAGGAAA

300





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| GTGCATGACT CACAGGGGAA | TOCAOCCAAA       | CCATGACCTC         | AGGAAGGAAA | GTGCATGACT | 360   |
|-----------------------|------------------|--------------------|------------|------------|-------|
| CACAGGOAGG AGCTGCTTAC | CCATOGAATG       | CAGCCAAACC         | ATGACCTCAG | TDAAADDAAD | 4 2 0 |
| GCATGACTOG GCAGCCAGCC | AGTOGCAGTT       | AATAGTGAAA         | CCCCGCCGAC | AGACATOTTT | 4 8 0 |
| TOCOAOCCTA OGAATCTTOO | CCTTOTCCCC       | AGTTAAACTO         | GACAAAGGCC | ATGGTTCTGC | 5 4 0 |
| GCCAGOCTGT CCTTCGAGCG | <b>dTdTTCCGC</b> | <b>o</b> tcctcctco | TATAGAAACT | CGGACCACTC | 6 0,0 |
| TOAGACGAAG OCTCGCGTCC | AGGCCAGCAC       | GAAGGAGGCT         | AAGTGGGAGG | OGTAGCOGTC | 660   |
| GTTGTCCACT AGGGGGTCCA | CTCGCTCCAG       | GOTGTGAAGA         | CACATGTCGC | CCTCTTCQQC | 720   |
| ATCAAGGAAG GTGATTGGTT | TATAGOTOTA       | GGCCAGACCG         | ootottcctg | TODDDDDDAA | 780   |
| ATAAAAGGG GTGGGGGCGC  | OTTCGTCCTC       | ACTCTCTTCC         | OCÀTCOCTOT | CTGCGAGGGC | 8 4 0 |
| CAGCIGATCA GCCTAGGCTT | TOCAMAMAGE       | TT                 |            | •,         | 872   |

#### ( 2 ) INFORMATION FOR SEQ ID NO.1:

## ( i ) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 663 base pairs (B) TYPH: mchio aid (C) STRANDEUNESS: single ( D ) TOPOLOGY: Saus

## ( i i ) MOLECULE TYPE: DNA (gene

## (x i ) SEQUENCE DESCRIPTION; SEQ ID NO:11:

| AAGCTTTTCT CATTAAGGGA | AGATTTCCCC | AGGCAGCTCT        | TTCAAGGCCT | AAAAGGTCCA | 6 0   |
|-----------------------|------------|-------------------|------------|------------|-------|
| TOAOCTCCAT OGATTCTTCC | CTOTTAAGAA | CTTTATCCAT        | TTTTGCAAAA | ATTGCAAAAG | 120   |
| AATAGGGATT TCCCCAAATA | GTTTTGCTAG | GCCTCAGAAA        | AAGCCTCCAC | ACCCTTACTA | 180   |
| CTTGAGAGAA AGGGTGGAGG | CAGAGGGGG  | CTCGGCCTTC        | TTATATATTA | *********  | 240   |
| GGCCACAGGG AGGAGCTGCT | TACCCATGGA | ATGCAGCCAA        | ACCATGACCT | CAGGAAGGAA | 300   |
| AGTGCATGAC TCACAGGGGA | ATGCAGCCAA | ACCATGACCT        | CAGGAAGGAA | AGTGCATGAC | 360   |
| TCACAGGAG GAGCTGCTTA  | CCCATGGAAT | GCAGCCAAAC        | CATGACCTCA | DAAADDAADD | 420   |
| TOCATOACTO OGCAGCCAGC | CAGTGGCAGT | TAATACAGGG        | TOTGAAGACA | CATGTCGCCC | 480   |
| TCTTCGGCAT CAAGGAAGGT | GAATTGGTTT | ATAGGTGTAG        | OCCACOTGAC | COGGTGTTCC | 5 4 0 |
| TOAAOGOGGO CTATAAAAGO | gaataaaac  | <b>GCGTTCGTCC</b> | TCACTCTCTT | CCGCATCGCT | 600   |
| GICTGCOAGG GCCAGTGATC | AGCCTAGGCT | TTGCAAAAAG        | CTT        |            | 643   |

### I claim:

- 1. A The recombinant human protein C molecule produced by inserting a vector comprising the DNA encoding human protein C into an adenovirus-transformed host cell then culturing said host cell under growth conditions suitable for production of said recombinant human protein C.
  - 2. The recombinant human protein C molecule of claim 1 wherein the adenovirus-transformed host cell is selected 55 from the group consisting of AV12 cells and human embryonic kidney 293 cells.
  - 3. The recombinant human protein C molecule of claim 2 wherein the adenovirus-transformed host cell is an AV12
  - 4. The recombinant human protein C molecule of claim 2 wherein the adenovirus transformed host cell is a human embryonic kidney 293 cell.